SEQUENCE LISTING

<110> Tsang et al.

<120> METHODS AND COMPOSITIONS FOR DETECTING LARVAL TAENIA SOLIUM

<130> 6395-62068

<140> 10/048,146
<141> 2000-08-03

<150> US 60/147,318
<151> 1999-08-03

<150> PCT/US00/21173
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<170> PatentIn version 3.1

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acttgtaact gaacaacctg taga atg cgt gcc tac att gtg ctt ctc gct   171
                           Met Arg Ala Tyr Ile Val Leu Leu Ala
                           1   5

ctc act gtt ttc gta gtg acg gtg tcg gcc gag tgg gtg ccc att tcg   219
Leu Thr Val Phe Val Val Thr Val Ser Ala Glu Trp Val Pro Ile Ser
10   15   20   25

agt gtc cac ata gcc tca tgc aaa agc cac tac atg ttc caa tta aaa   267
Ser Val His Ile Ala Ser Cys Lys Ser His Tyr Met Phe Gln Leu Lys
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cgc ttt ttt gcc ttt agg aaa aac aaa ccg aaa gat gtt gca aat agt   315
Arg Phe Phe Ala Phe Arg Lys Asn Lys Pro Lys Asp Val Ala Asn Ser
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acg aaa aaa ggg ata gaa tat gtc cac gaa ttc ttc cac gaa gac ccg   363
Thr Lys Lys Gly Ile Glu Tyr Val His Glu Phe Phe His Glu Asp Pro
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att ggt aaa caa att gct caa ctc gca aag gaa tgg aag gaa gca atg        411
Ile Gly Lys Gln Ile Ala Gln Leu Ala Lys Glu Trp Lys Glu Ala Met
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ttg gaa ggt agg ttt tgg tgt ttt ctg tca gaa gaa aat tat cta ttc        459
Leu Glu Gly Arg Phe Trp Cys Phe Leu Ser Glu Glu Asn Tyr Leu Phe
90                  95                  100                 105

att cat cta gac aaa ggc aaa ata cgg acg tca ctg gtt gag cac tgc        507
Ile His Leu Asp Lys Gly Lys Ile Arg Thr Ser Leu Val Glu His Cys
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aaa ggt cct aag aaa aaa act gct taacttgtca actttcatgc gttcttctct       561
Lys Gly Pro Lys Lys Lys Thr Ala
                125

tcactaataa atgctcatta ataagaaagc tgcctttgc aagatcaacg agggccatag        621

actgtgaggg ttatagccta aggttatggg gtgaaatgag ataggaattg agcatttgag        681

aagttactaa tttaaattga aagccgcatt cttctgcaa ttgacgtgtg atggttagcg         741

aaaccaagtg aagcacgacc tcttgagtcg tttcaacagc cgccagtggt ttcaccagtg        801

gcttcaccag tgggtagact ggtttgtcac acatgcgagg tacggtcaga gggctaacag        861

gtgtggtgga ggggccaaca cgtgtaagac aagcagttcc ccttctctgt cgtgaggcac        921

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gagggtagga tgagtgtaca gaaaacagcg aggcaacgaa tctactggca tggccctgat       1581

gccacccgc ccagctaggg tagtttggcc acctcagtcc ttaatcgaat gcggcagtca        1641

gaacaaacaa agtattacat agccacactc ttcttttgag cgtcgtcctc gacgctcctt       1701

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aacgatttca ctttgatgct ctcaccatca attctcttgt atgtgccacc accttaaacc   2061

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Lys Ser His Tyr Met Phe Gln Leu Lys Arg Phe Phe Ala Phe Arg Lys
            35                  40                  45


Asn Lys Pro Lys Asp Val Ala Asn Ser Thr Lys Lys Gly Ile Glu Tyr
        50                  55                  60


Val His Glu Phe Phe His Glu Asp Pro Ile Gly Lys Gln Ile Ala Gln
65                  70                  75                  80


Leu Ala Lys Glu Trp Lys Glu Ala Met Leu Glu Gly Arg Phe Trp Cys
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Phe Leu Ser Glu Glu Asn Tyr Leu Phe Ile His Leu Asp Lys Gly Lys
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gca aat agt act aag aaa gag ata gaa tat atc cac aat tgg ttt ttc   95
Ala Asn Ser Thr Lys Lys Glu Ile Glu Tyr Ile His Asn Trp Phe Phe
                20   25   30

cat gat gac ccg att gga aaa caa att gct caa ctc gca aag gac tgg   143
His Asp Asp Pro Ile Gly Lys Gln Ile Ala Gln Leu Ala Lys Asp Trp
            35   40   45

aat gaa aca gtg cag gaa gcc aaa ggc aaa ttt tgg gcg tca ctg gct   191
Asn Glu Thr Val Gln Glu Ala Lys Gly Lys Phe Trp Ala Ser Leu Ala
        50   55   60

gag tac tgc aga ggt ctg aag aac aaa act gct taacttgtca actttcatgc   244
Glu Tyr Cys Arg Gly Leu Lys Asn Lys Thr Ala
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Asp Asp Pro Ile Gly Lys Gln Ile Ala Gln Leu Ala Lys Asp Trp Asn
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gta aag aat att aag aaa ggg atg gaa gtt gtc tac aaa ttt ttc tac       95
Val Lys Asn Ile Lys Lys Gly Met Glu Val Val Tyr Lys Phe Phe Tyr
                20                  25                  30

gaa gac ccg ttg gga aag aaa ata gct caa ctc gca aag gac tgg aag      143
Glu Asp Pro Leu Gly Lys Lys Ile Ala Gln Leu Ala Lys Asp Trp Lys
                35                  40                  45

gaa gca atg ttg gaa gcc aga agc aaa gtg cgg gcg tca ctg gct gag      191
Glu Ala Met Leu Glu Ala Arg Ser Lys Val Arg Ala Ser Leu Ala Glu
            50                  55                  60

tac atc aga ggt ctc aag aac gaa gct gct taacttgtca actttcatgc        241
Tyr Ile Arg Gly Leu Lys Asn Glu Ala Ala
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5

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Ala Met Leu Glu Ala Arg Ser Lys Val Arg Ala Ser Leu Ala Glu Tyr
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Ile Arg Gly Leu Lys Asn Glu Ala Ala
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Ile Ala Gln Leu Ala Lys
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<223> Asparagine at position 21 is an amino acid insertion

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Tyr Ile Trp His Asn Phe Phe Phe
            20


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<220>
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<223>  Amino acid at position 12 may also be aspartic acid


<220>
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<220>
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<223>  Tryptophan at position 8 is an amino acid insertion


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Gly Ile Glu Tyr Val His Glu Trp Phe Phe His Glu Asp
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